

1600

RAW SEQUENCE LISTING

DATE: 12/07/2001

PATENT APPLICATION: US/09/665,350

TIME: 14:31:39

Input Set : N:\EBONY'S\ES.txt

Output Set: N:\CRF3\12072001\I665350.raw

p.5

ENTERED

3 <110> APPLICANT: Genentech, Inc.
4 Ashkenazi, Avi
5 Botstein, David
6 Desnoyers, Luc
7 Eaton, Dan L.
8 Ferrara, Napoleone
9 Filvaroff, Ellen
10 Fong, Sherman
11 Gao, Wei-Qiang
12 Gerber, Hanspeter
13 Gerritsen, Mary E.
14 Goddard, A.
15 Godowski, Paul J.
16 Grimaldi, Christopher J.
17 Gurney, Austin L.
18 Hillan, Kenneth, J.
19 Kljavin, Ivar J.
20 Mather, Jennie P.
21 Pan, James
22 Paoni, Nicholas F.
23 Roy, Margaret Ann
24 Stewart, Timothy A.
25 Tumas, Daniel
26 Williams, P. Mickey
27 Wood, William, I.
29 <120> TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
30 Acids Encoding the Same
32 <130> FILE REFERENCE: 10466-14
34 <140> CURRENT APPLICATION NUMBER: 09/665,350
35 <141> CURRENT FILING DATE: 2000-09-18
37 <150> PRIOR APPLICATION NUMBER: PCT/US00/04414
38 <151> PRIOR FILING DATE: 2000-02-22
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41 <151> PRIOR FILING DATE: 1999-07-07
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47 <151> PRIOR FILING DATE: 1999-07-28
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50 <151> PRIOR FILING DATE: 1999-09-08
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53 <151> PRIOR FILING DATE: 1999-09-13
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155 aaaattgacc attgtaggta atcaggagga aaaaaaaaaa aaaaaaaaaa 1450
157 aaagggcggc cgcgactcta gagtcgacct gcagaagctt ggccgccatg 1500
159 gcccaacttg tttattgcag cttataatgg ttacaaataa agcaatagca 1550
161 tcacaaattt cacaaataaa gcattttttt cactgcattc tagttgtggt 1600
163 ttgtccaaac tcatcaatgt atcttatcat gtctggatcg ggaattaatt 1650
165 cggcgagca ccatggcctg aaataacctc tgaaagagga acttggttag 1700
167 gtaccttctg aggcggaaag aaccagctgt ggaatgtgtg tcagttaggg 1750
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171 ctcaattagt cagcaaccca gtttt 1825
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185 Cys His Arg Cys Arg Gly Leu Val Asp Lys Phe Asn Gln Gly Met
186 35 40 45
188 Val Asp Thr Ala Lys Lys Asn Phe Gly Gly Gly Asn Thr Ala Trp
189 50 55 60
191 Glu Glu Lys Thr Leu Ser Lys Tyr Glu Ser Ser Glu Ile Arg Leu
192 65 70 75
194 Leu Glu Ile Leu Glu Gly Leu Cys Glu Ser Ser Asp Phe Glu Cys
195 80 85 90
197 Asn Gln Met Leu Glu Ala Gln Glu Glu His Leu Glu Ala Trp Trp
198 95 100 105
200 Leu Gln Leu Lys Ser Glu Tyr Pro Asp Leu Phe Glu Trp Phe Cys
201 110 115 120
203 Val Lys Thr Leu Lys Val Cys Cys Ser Pro Gly Thr Tyr Gly Pro
204 125 130 135
206 Asp Cys Leu Ala Cys Gln Gly Gly Ser Gln Arg Pro Cys Ser Gly
207 140 145 150
209 Asn Gly His Cys Ser Gly Asp Gly Ser Arg Gln Gly Asp Gly Ser
210 155 160 165
212 Cys Arg Cys His Met Gly Tyr Gln Gly Pro Leu Cys Thr Asp Cys
213 170 175 180
216 Met Asp Gly Tyr Phe Ser Ser Leu Arg Asn Glu Thr His Ser Ile
217 185 190 195
219 Cys Thr Ala Cys Asp Glu Ser Cys Lys Thr Cys Ser Gly Leu Thr
220 200 205 210
222 Asn Arg Asp Cys Gly Glu Cys Glu Val Gly Trp Val Leu Asp Glu
223 215 220 225
225 Gly Ala Cys Val Asp Val Asp Glu Cys Ala Ala Glu Pro Pro Pro
226 230 235 240
228 Cys Ser Ala Ala Gln Phe Cys Lys Asn Ala Asn Gly Ser Tyr Thr

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Input Set : N:\EBONY'S\ES.txt

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232          260          265          270
234 Pro Gly Asn Cys Lys Glu Cys Ile Ser Gly Tyr Ala Arg Glu His
235          275          280          285
237 Gly Gln Cys Ala Asp Val Asp Glu Cys Ser Leu Ala Glu Lys Thr
238          290          295          300
240 Cys Val Arg Lys Asn Glu Asn Cys Tyr Asn Thr Pro Gly Ser Tyr
241          305          310          315
243 Val Cys Val Cys Pro Asp Gly Phe Glu Glu Thr Glu Asp Ala Cys
244          320          325          330
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247          335          340          345
249 Gln Leu Pro Ser Arg Glu Asp Leu
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252 <210> SEQ ID NO: 3
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262 cgcccagccg tctaaacggg aacagccctg gctgaggag ctgcagcgca 150
264 gcagagtatc tgacggcgcc aggttgcgta ggtgcggcac gaggagtttt 200
266 cccggcagcg aggaggtcct gaggcagcatg gcccgaggga gcgccttccc 250
268 tgccgcccgcg ctctggctct ggagcatcct cctgtgcctg ctggcactgc 300
270 gggcgagggc cgggcccgcg caggaggaga gcctgtacct atggatcgat 350
272 gctcaccagg caagagtact cataggattt gaagaagata tcctgattgt 400
274 ttcagagggg aaaatggcac cttttacaca tgatttcaga aaagcgcaac 450
276 agagaatgcc agctattcct gtcaatatcc attccatgaa ttttacctgg 500
278 caagctgcag ggcaggcaga atacttctat gaattcctgt ccttgcgctc 550
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283 gaacagtgcc tcacaaggca tcagttgttc aagttggttt cccatgtctt 650
285 ggaaaacagg atggggtggc agcatttgaa gtggatgtga ttgttatgaa 700
287 ttctgaaggc aacaccattc tccaaacacc tcaaaatgct atcttcttta 750
289 aaacatgtca acaagctgag tgcccaggcg ggtgccgaaa tggaggcttt 800
291 tgtaatgaaa gacgcatctg cgagtgtcct gatgggttcc acggacctca 850
293 ctgtgagaaa gccctttgta cccacgatg tatgaatggg ggactttgtg 900
295 tgactcctgg tttctgcac tgcccacctg gattctatgg agtgaactgt 950
297 gacaaagcaa actgctcaac cacctgcttt aatggaggga cctgtttcta 1000
299 ccctggaaaa tgtatttgcc ctccaggact agagggagag cagtgtgaaa 1050
301 tcagcaaata cccacaaccc tgtcgaaatg gaggtaaatg catttgtaaa 1100
303 agcaaatagt agtgttccaa aggttaccag ggagacctct gttcaaagcc 1150
305 tgtctgcgag cctggctgtg gtgcacatgg aacctgccat gaaccaaca 1200
307 aatgccaatg tcaagaagg tggcatggaa gacactgcaa taaaaggtac 1250
309 gaagccagcc tcatacatgc cctgaggcca gcaggcgccc agctcaggca 1300
311 gcacacgcct tcacttaaaa aggccgagga gcggcgggat ccacctgaat 1350
313 ccaattacat ctggtgaact ccgacatctg aaacgtttta agttacacca 1400
315 agttcatagc ctttgttaac ctttcatgtg ttgaatgttc aaataatgtt 1450

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Output Set: N:\CRF3\12072001\I665350.raw

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319 actgagctga tatttactct tccttttaag ttttctaagt acgtctgtag 1550
321 catgatggta tagattttct tgtttcagtg ctttgggaca gattttatat 1600
323 tatgtcaatt gatcagggtta aaattttcag tgtgtagttg gcagatattt 1650
325 tcaaaattac aatgcattta tgggtgtctgg gggcagggga acatcagaaa 1700
327 ggttaaattg ggcaaaaatg cgtaagtcac aagaatttgg atggtgcagt 1750
329 taatgttgaa gttacagcat ttcagatttt attgtcagat atttagatgt 1800
331 ttgttacatt tttaaaaatt gctcttaatt tttaaactct caatacaata 1850
333 tattttgacc ttaccattat tccagagatt cagtattaaa aaaaaaaaaa 1900
335 ttacactgtg gtagtggcat ttaaacaata taatatattc taaacacaat 1950
337 gaaataggga atataatgta tgaacttttt gcattggctt gaagcaatat 2000
339 aatataattgt aaacaaaaca cagctcttac ctaataaaca ttttatactg 2050
341 tttgtatgta taaaataaag gtgctgcttt agttttttgg aaaaaaaaaa 2100
343 aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa gggcgggcgc gactctagag 2150
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360 20 25 30
362 Pro Gln Glu Glu Ser Leu Tyr Leu Trp Ile Asp Ala His Gln Ala
363 35 40 45
365 Arg Val Leu Ile Gly Phe Glu Glu Asp Ile Leu Ile Val Ser Glu
366 50 55 60
368 Gly Lys Met Ala Pro Phe Thr His Asp Phe Arg Lys Ala Gln Gln
369 65 70 75
371 Arg Met Pro Ala Ile Pro Val Asn Ile His Ser Met Asn Phe Thr
372 80 85 90
374 Trp Gln Ala Ala Gly Gln Ala Glu Tyr Phe Tyr Glu Phe Leu Ser
375 95 100 105
377 Leu Arg Ser Leu Asp Lys Gly Ile Met Ala Asp Pro Thr Val Asn
378 110 115 120
380 Val Pro Leu Leu Gly Thr Val Pro His Lys Ala Ser Val Val Gln
381 125 130 135
383 Val Gly Phe Pro Cys Leu Gly Lys Gln Asp Gly Val Ala Ala Phe
384 140 145 150
386 Glu Val Asp Val Ile Val Met Asn Ser Glu Gly Asn Thr Ile Leu
387 155 160 165
389 Gln Thr Pro Gln Asn Ala Ile Phe Phe Lys Thr Cys Gln Gln Ala
390 170 175 180
392 Glu Cys Pro Gly Gly Cys Arg Asn Gly Gly Phe Cys Asn Glu Arg
393 185 190 195
395 Arg Ile Cys Glu Cys Pro Asp Gly Phe His Gly Pro His Cys Glu
396 200 205 210

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Use of n and/or Xaa has been detected in the Sequence Listing.
 Review the Sequence Listing to insure a corresponding
 explanation is presented in the <220> to <223> fields of
 each sequence using n or Xaa.

VERIFICATION SUMMARY

DATE: 12/07/2001

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Output Set: N:\CRF3\12072001\I665350.raw

L:651 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:653 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:655 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:657 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:978 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26
L:2194 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50
L:4666 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:113
L:5251 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:131
L:6947 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:174
L:7127 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:175
L:8523 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:206
L:8525 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:206